

FIG 1C

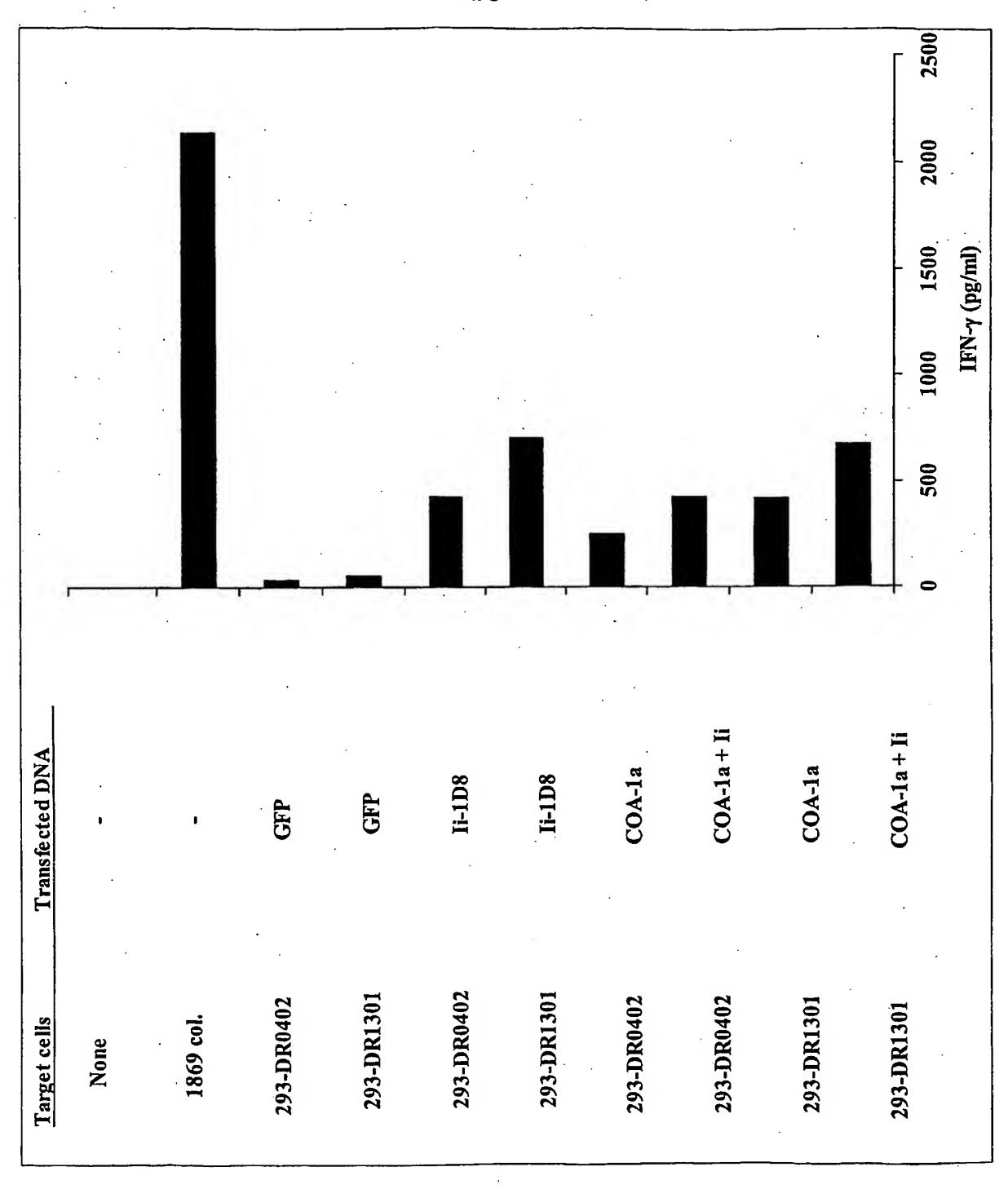


FIG. 2

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MAFMTRKLWD LEQQVKAQTD EILSKDQKIA ALEDLVQTLR PHPAEATLQR QEELETMCVQ 60

LQRQVREMER FLSDYGLQWV GEPMDQEDSE SKTVSEHGER DWMTAKKFWK PGDSLAPPEV 120

DFDRLLASLQ DLSELVVEGD TQVTPVPGGA RLRTLEPIPL KLYRNGIMMF DGPFQPFYDP 180

STQRCLRDIL DGFFPSELQR LYPNGVPFKV SDLRNQVYLE DGLDPFPGEG RVVGRQRMHK 240

ALDRVEEHPG SRMTAEKFLN RLPKFVIRQG EVIDIRGPIR DTLQNCCPLP ARIQEIVVET 300

PTLAAERERS QESPNTPAPP LSMLRIKSEN GEQAFLLMMQ PDNTIGDVRA LLAQARVMDA 360

SAFEIFSTFP PTLYQDDTLT LQAAGLVPKA ALLLRARRAP KSSLKFSPGP CPGPGPGPSP 420

GPGPGSSPCP GPSPSPQ 437

FIG. 3

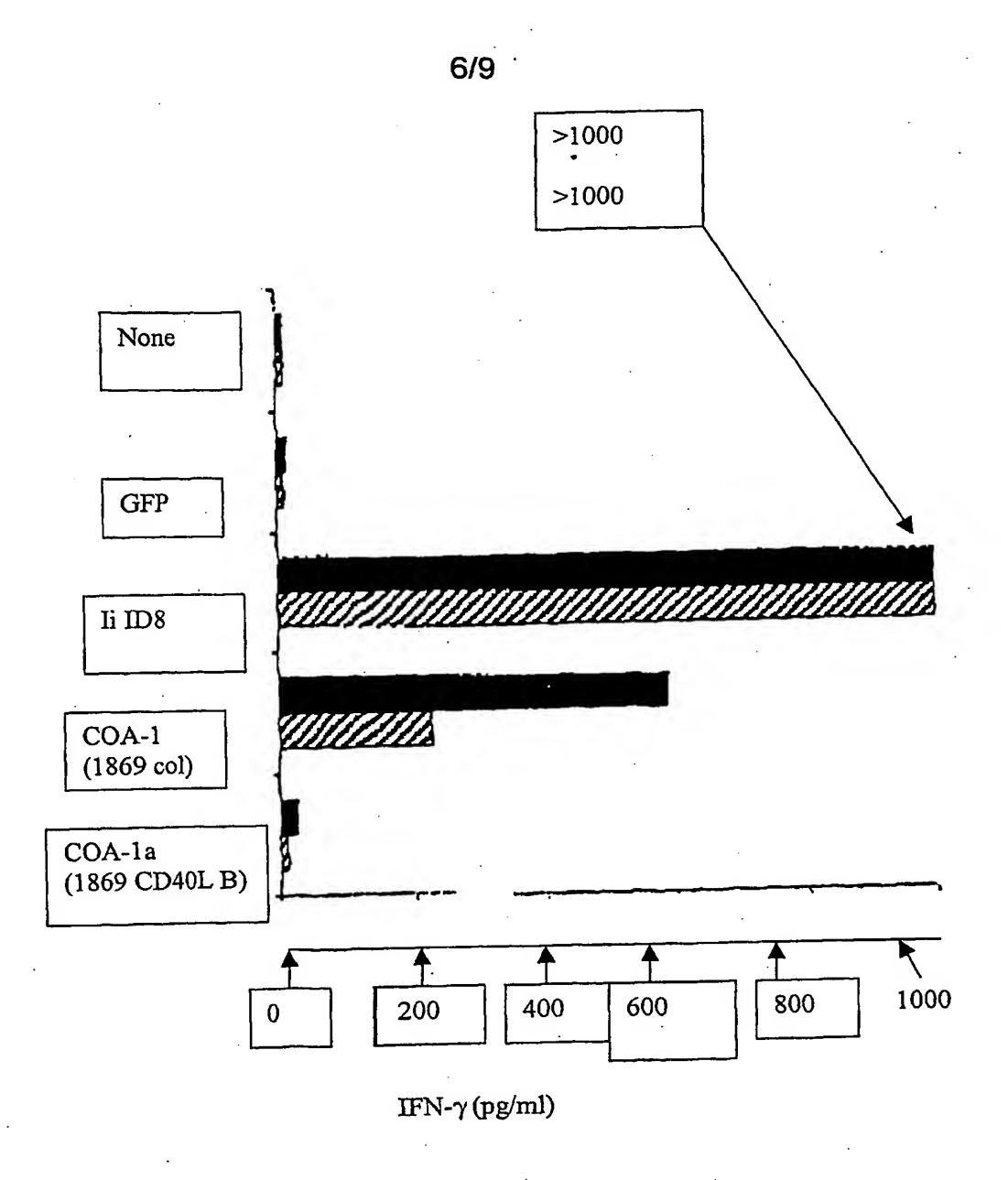


FIG. 4

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	cgctgcggga cggctagcgg ccctgcgtgg aggcgaggaa tccgcatcta tggagatgtc													60			
	cctg	cato	:CC a	tgac	tegg	ja go				•			agg a Arg I		_	_	111
•													ata Ile		_		159
	_												acc Thr			cca Pro	207
													ctg Leu		_	_	255
	tgt Cys	gtg Val	cag Gln 60	ctg Leu	cag Gln	cgg Arg	cag Gln	gtc Val 65	agg Arg	gag Glu	atg Met	gag Glu	cgg Arg 70	ttc Phe	ctc Leu	agt Ser	303
-													cag Gln				351
								_		_			tgg Trp	_			399
	aag Lys	aag Lys	ttc Phe	tgg Trp	aag Lys 110	cca Pro	elà aaa	gac Asp	tca Ser	ttg Leu 115	Ala	ccc Pro	cct Pro	gag Glu	gtg Val 120	gac Asp	447
													gag Glu		Val		495
	gag Glu	ggt Gly	gac Asp 140	acc Thr	caa Gln	gtg Väl	aca Thr	cca Pro 145	gtg Val	ccc Pro	Gly	gjy aaa	gca Ala 150	cgg Arg	ctg Leu		543
	acc Thr	ctc Leu 155	gag Glu	ccc Pro	atc Ile	ccġ Pro	Leu	Lys	Leu	Tyr	Arg	Asn	ggc	atc Ile	atg Met	atg Met	591
	ttc Phe 170	gac Asp	gjå aaa	ccc Pro	ttc Phe	cag Gln 175	ccc Pro	ttc Phe	tac Tyr	gat Asp	ccc Pro 180	Ser	aca Thr	cag Gln	cgc	tgc Cys 185	639
	ctc Leu	cga Arg	gac Asp	ata Ile	ttg Leu 190	gat Asp	ggc Gly	ttc Phe	ttt Phe	ccc Pro 195	tca Ser	gag Glu	ctc Leu	cag Gln	cga Arg 200	ctg Leu	687
													cgc Arg				735

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		_	gat Asp					_					_	_	_	783
_	_		Cgg Arg		_		_									831
	Ser		atg Met											_		879
_			caa Gln	,	=	_		-							-	927
	_		aac Asn 285	_	_											.975
		_	bio CCC		_	_	- -		•							1023
			ccg Pro													1071
			cag Gln						Met							1119
			cga Arg												Ser	1167
			atc Ile 365						Pro							1215
aca Thr	ctc Leu	acg Thr 380	ctg Leu	cag Gln	gct Ala	gca Ala	ggc Gly 385	ctt Leu	gtg Val	ccc	aaa Lys	gça Ala 390	gca Ala	ctg Leu	ctg Leu	1263
		٠					Cyte	osine	at po	ositio	n 128	30				

ctg cgg gca cgc cga g ${f C}$ c ccg aag tcc agc ctg aaa ttc agt cct ggt 1311

Leu Arg Ala Arg Arg **Ala** Pro Lys Ser Ser Leu Lys Phe Ser Pro Gly
395 400 405

Alanine at position 399

ccc tgt ccc ggt ccc ggt ccc ggc ccc agt ccc ggt ccc ggt ccc ggc 1359
Pro Cys Pro Gly Pro Gly Pro Gly Pro Ser Pro Gly Pro Gly Pro Gly
410 425

FIG. 5 cont.

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tcc Ser	agt Ser	ccc Pro	tgt Cys	ccc Pro 430	gga Gly	ccc Pro	agt Ser	ccc Pro	agc Ser 435	Pro	caa Gln	ta	aag	cacc	CC .	٠	1405	
acco	iccto	3													-		1413	

FIG. 5 cont.